

STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING

ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: _____

Source: _____

Date Processed by STIC: _____

10/550,608
LFWP
11/14/2006

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

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- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.4.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
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Revised 01/10/06



IFWP

RAW SEQUENCE LISTING

DATE: 11/14/2006

PATENT APPLICATION: US/10/550,608

TIME: 08:51:21

Input Set : N:\RJAVED\10550608.txt

Output Set: N:\CRF4\11142006\J550608.raw

4 <110> APPLICANT: Progenika Biopharma, S.A.

6 <120> TITLE OF INVENTION: In vitro method to detect bladder transitional cell carcinoma

C--> 8 <140> CURRENT APPLICATION NUMBER: US/10/550,608

C--> 8 <141> CURRENT FILING DATE: 2005-09-26.

W--> 0 <130> FILE REFERENCE:

8 <160> NUMBER OF SEQ ID NOS: 18

10 <170> SOFTWARE: PatentIn version 3.1

Does Not Only
Corrected Sequence Needed
Cp9-1)

ERRORED SEQUENCES

282 <210> SEQ ID NO: 18

283 <211> LENGTH: 25

284 <212> TYPE: DNA

285 <213> ORGANISM: Artificial sequence

W--> 287 <220> FEATURE: synthetic DNA

288 <223> OTHER INFORMATION: probe sequence of the 31805_at probe set of Affymetrix, the position of said probe in the mRNA sequence of the fgfr3 gene being

289 3720

293 <400> SEQUENCE: 18

294 gtggccagag gtgtcaccca aaccg 25

E--> 299 1

Pls delete

DO not Insert any
Information in this
mandatory Identific

This type of Error
Occur subsequently
throughout the Sequence.

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/550,608

DATE: 11/14/2006

TIME: 08:51:22

Input Set : N:\RJAVED\10550608.txt

Output Set: N:\CRF4\11142006\J550608.raw

L:8 M:270 C: Current Application Number differs, Replaced Current Application No
L:8 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:0 M:201 W: Mandatory field data missing, <130> FILE REFERENCE
L:17 M:256 W: Invalid Numeric Header Field, <220> has non-blank data
L:32 M:256 W: Invalid Numeric Header Field, <220> has non-blank data
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L:299 M:254 E: No. of Bases conflict, this line has no nucleotides.